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WHAT IS CLAIMED IS:

1	1. A method for reducing a condition associated with fetal alcohol
2	syndrome in a subject who is exposed to alcohol in utero, the method comprising
3	administering to the subject an ADNF polypeptide in an amount sufficient to reduce the
4	condition associated with fetal alcohol syndrome.
1	2. The method of claim 1, wherein the ADNF polypeptide is a
2	member selected from the group consisting of:
3	(a) an ADNF I polypeptide comprising an active core site having the
4	following amino acid sequence:
5	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1);
6	(b) an ADNF III polypeptide comprising an active core site having the
7	following amino acid sequence:
8	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2); and
9	(c) a mixture of the ADNF I polypeptide of part (a) and the ADNF III
10	polypeptide of part (b).
1	3. The method of claim 1, wherein the ADNF polypeptide is a
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2	member selected from the group consisting of a full length ADNF I polypeptide, a full
3	length ADNF III polypeptide, and a mixture of a full length ADNF I polypeptide and a
4	full length ADNF III polypeptide.
1	4. The method of claim 1, wherein the ADNF polypeptide is an
2	ADNF I polypeptide.
1	5. The method of claim 4, wherein the ADNF I polypeptide is Ser-
2	Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1).
1	6. The method of claim 4, wherein the ADNF I polypeptide is
2	selected from the group consisting of:
3	Val-Leu-Gly-Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:14);
4	Val-Clu-Gly-Gly-Gly-Ser-Ala-Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-
- 5	Ala (SEQ ID NO:15);
6	Leu-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:16);
U	Lea-Oly-Oly-Oly- bel-Ala-Lea-Lea-Alg-bel-He-f 10-Ala (bbQ ib 110.10);

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7	Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:17);
8	Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:18); and
9	Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:19).
1	7. The method of claim 4, wherein the ADNF I polypeptide
2	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
3	of the active core site.
1	8. The method of claim 1, wherein the ADNF polypeptide is an
2	ADNF III polypeptide.
1	9. The method of claim 8, wherein the ADNF III polypeptide is Asn-
2	Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	10. The method of claim 8, wherein the ADNF III polypeptide is
2	selected from the group consisting of:
3	Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:20);
4	Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID NO:21);
5	Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID
6	NO:22); and
7	Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser
8	(SEQ ID NO:23).
1	11. The method of claim 8, wherein the ADNF III polypeptide
2	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
3	of the active core site.
1	12. The method of claim 1, wherein the ADNF polypeptide is a
2	mixture of an ADNF I polypeptide of part (a) and an ADNF III polypeptide of part (b).
1	13. The method of claim 12, wherein the ADNF I polypeptide is Ser-
2	Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1), and wherein the ADNF III
3	polypeptide is Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	14. The method of claim 12, wherein the ADNF I polypeptide is
2	selected from the group consisting of:
3	Val-Leu-Gly-Gly-Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:14):

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4	Val-Glu-Glu-Gly-Ile-Val-Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-
5	Ala (SEQ ID NO:15);
6	Leu-Gly-Gly-Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:16);
7	Gly-Gly-Gr-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:17);
8	Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:18);
9	Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:19); and
10	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1); and wherein the ADNF II
11	polypeptide is selected from the group consisting of:
12	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2);
13	Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:20);
14	Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID NO:21);
15	Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID
16	NO:22); and
17	Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser
18	(SEQ ID NO:23).
1	15. The method of claim 12, wherein the ADNF I polypeptide
2	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
3	of the active core site of the ADNF I polypeptide, and wherein the ADNF III polypeptide
4	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
5	of the active core site of the ADNF III polypeptide.
1	16. The method of claim 1, wherein at least one of the ADNF
2	polypeptide is encoded by a nucleic acid which is administered to the subject.
1	17. The method of claim 1, wherein the condition is decreased body
2	weight of the subject.
1	18. The method of claim 1, wherein the condition is decreased brain
2	weight of the subject.
1	19. The method of claim 1, wherein the condition is a decreased level
2	of VIP mRNA or protein of the subject.
1	20. The method of claim 1, wherein the condition is decreased viability
2	of the subject in utero.





1	21. The method of claim 1, wherein the condition is decreased
2	learning.
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1	A method for reducing neuronal cell death, the method comprising
2	contacting a neuronal cell with a mixture of an ADNF I polypeptide and an ADNF III
3	polypeptide in an amount sufficient to reduce neuronal cell death,
4	wherein the ADNF I polypeptide comprises an active core site having the
5	following amino acid sequence:
6	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1); and
7	wherein the ADNF III polypeptide comprises an active core site having the
8	following amino acid sequence:
9	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	23. The method of claim 22, wherein the ADNF I polypeptide is a full
2	length ADNF I polypeptide and the ADNF III polypeptide is a full length ADNF III
3	polypeptide.
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1	24. The method of claim 22, wherein the ADNF I polypeptide is Ser-
2	Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1), and wherein the ADNF III
3	polypeptide is Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	25. The method of claim 22, wherein the ADNF I polypeptide is
2	selected from the group consisting of:
3	Val-Leu-Gly-Gly-Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:14);
4	Val-Glu-Glu-Gly-Ile-Val-Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-
5	Ala (SEQ ID NO:15);
6	Leu-Gly-Gly-Gly- Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:16);
7	Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:17);
8	Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:18);
9	Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:19); and
10	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1); and wherein the ADNF III
11	polypeptide is selected from the group consisting of:
12	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2);
13	Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:20);
14	Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID NO:21);

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15	Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID
16	NO:22); and
17	Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser
18	(SEQ ID NO:23).
1	26. The method of claim 22, wherein the ADNF I polypeptide
2	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
3	of the active core site of the ADNF I polypeptide, and wherein the ADNF III polypeptide
4	comprises up to about 20 amino acids at at least one of the N-terminus and the C-terminus
5	of the active core site of the ADNF III polypeptide.
1	27. The method of claim 22, wherein at least one of the ADNF
2	polypeptide is encoded by a nucleic acid.
1	28. A pharmaceutical composition comprising a pharmaceutically
2	acceptable excipient and a mixture of an ADNF I polypeptide and an ADNF III
3	polypeptide, wherein the ADNF I polypeptide comprises an active core site having the
4	following amino acid sequence:
5	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1); and
6	wherein the ADNF III polypeptide comprises an active core site having the following
7	amino acid sequence:
8	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	29. The pharmaceutical composition of claim 28, wherein the ADNF I
2	polypeptide is a full length ADNF I polypeptide and the ADNF III polypeptide is a full
3	length ADNF III polypeptide.
1	30. The pharmaceutical composition of claim 28, wherein the ADNF I
2	polypeptide is Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1), and wherein the
3	ADNF III polypeptide is Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2).
1	The pharmaceutical composition of claim 28, wherein the ADNF I
2	polypeptide is selected from the group consisting of:
3	Val-Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:14);
4	Val-Glu-Glu-Gly-Ile-Val-Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-
5	Ala (SEO ID NO:15);

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6	Leu-Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:16);
7	Gly-Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:17);
8	Gly-Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:18)
9	Gly-Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:19); and
10	Ser-Ala-Leu-Leu-Arg-Ser-Ile-Pro-Ala (SEQ ID NO:1); and wherein the ADNF III
11	polypeptide is selected from the group consisting of:
12	Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:2)
13	Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:20);
14	Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID NO:21);
15	Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser (SEQ ID
16	NO:22); and
17	Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-Gln-Ser
18	(SEQ ID NO:23).
1	32. The pharmaceutical composition of claim 28, wherein the ADNF I
2	polypeptide comprises up to about 20 amino acids at at least one of the N-terminus and
3	the C-terminus of the active core site of the ADNF I polypeptide, and wherein the ADNF
4	III polypeptide comprises up to about 20 amino acids at at least one of the N-terminus and
5	the C-terminus of the active core site of the ADNF III polypeptide.
1	33. The pharmaceutical composition of claim 28, wherein at least one
2	of the ADNF polypeptide is encoded by a nucleic acid.